STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _10/587,370_
Source: _1 FWP_
Date Processed by STIC: _8/7/06_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
       http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (**<http://www.uspto.gov/ebc/efs/downloads/documents.htm>** , **EFS Submission User Manual** - ePAVE)
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):**
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06

Raw Sequence Listing Error Summary

ERROR DETECTED   **SUGGESTED CORRECTION**   SERIAL NUMBER: _10/587,370_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1____Wrapped Nucleics   The number/text at the end of each line "wrapped" down to the next line. This may occur if your file
      Wrapped Aminos   was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will
                          prevent "wrapping."

2____Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

3____Misaligned Amino   The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers;
      Numbering   use **space characters**, instead.

4____Non-ASCII   The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please
                          ensure your subsequent submission is saved in ASCII text.**

5____Variable Length   Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules,
                          each n or Xaa can only represent a single residue.** Please present the **maximum** number of each
                          residue having variable length and indicate in the <220>-<223> section that some may be missing.

6____PatentIn 2.0   A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
      "bug"   sequences(s) _____. Normally, PatentIn would automatically generate this section from the
                          previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to
                          the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for
                          Artificial or Unknown sequences.**

7____Skipped Sequences   Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence:
      (OLD RULES)   (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                          (i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
                          (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
                          This sequence is intentionally skipped
                          Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

8____Skipped Sequences   Sequence(s) _____ missing. If **intentional**, please insert the following lines for **each** skipped sequence.
      (NEW RULES)   <210> sequence id number
                          <400> sequence id number
                          000

9____Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
      (NEW RULES)   Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
                          In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.

10____Invalid <213>   Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or
       Response   scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or
                          is Artificial Sequence. (see item 11 below)

11____Use of <220>   Sequence(s) ____ missing the <220> "Feature" and associated numeric identifiers and responses. Use
                          of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown."
                          Please explain source of genetic material in <220> to <223> section or use "chemically synthesized" as
                          explanation. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32), also Sec. 1.823 of
                          Sequence Rules

12____PatentIn 2.0   Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
       "bug"   resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence
                          listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13____Misuse of n/Xaa   "n" can **only** represent a single <u>nucleotide</u>; "Xaa" can **only** represent a single <u>amino acid</u>

AMC – STIC Systems Branch – 03/02/06

RAW SEQUENCE LISTING   DATE: 08/07/2006
PATENT APPLICATION: US/10/587,370   TIME: 09:35:48

Input Set : **A:\11669.0237USWO-Sequence Listing.txt**
Output Set: **N:\CRF4\08072006\J587370.raw**

```
     3 <110> APPLICANT: Genentech, Inc.
     5 <120> TITLE OF INVENTION: BCMA POLYPEPTIDES AND USES THEREOF
     7 <130> FILE REFERENCE: 11669.0237USWO
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/587,370
    10 <141> CURRENT FILING DATE: 2006-07-26
    12 <150> PRIOR APPLICATION NUMBER: PCT/US2004/025247
    13 <151> PRIOR FILING DATE: 2004-08-04
    15 <150> PRIOR APPLICATION NUMBER: US 60/540,271
    16 <151> PRIOR FILING DATE: 2004-01-29
    18 <160> NUMBER OF SEQ ID NOS: 33
    20 <170> SOFTWARE: PatentIn version 3.3
    22 <210> SEQ ID NO: 1
    23 <211> LENGTH: 34
    24 <212> TYPE: PRT
    25 <213> ORGANISM: Artificial Sequence
    27 <220> FEATURE:
    28 <223> OTHER INFORMATION: Formula I
    31 <220> FEATURE:
    32 <221> NAME/KEY: MISC_FEATURE
    33 <222> LOCATION: (2)..(5)
    34 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
    36 <220> FEATURE:
    37 <221> NAME/KEY: MISC_FEATURE
    38 <222> LOCATION: (7)..(7)
    39 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
    41 <220> FEATURE:
    42 <221> NAME/KEY: MISC_FEATURE
    43 <222> LOCATION: (9)..(9)
    44 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
    46 <220> FEATURE:
    47 <221> NAME/KEY: MISC_FEATURE
    48 <222> LOCATION: (11)..(11)
    49 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
    51 <220> FEATURE:
    52 <221> NAME/KEY: MISC_FEATURE
    53 <222> LOCATION: (12)..(13)
    54 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
    56 <220> FEATURE:
    57 <221> NAME/KEY: MISC_FEATURE
    58 <222> LOCATION: (16)..(16)
    59 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
    61 <220> FEATURE:
    62 <221> NAME/KEY: MISC_FEATURE
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[handwritten annotations:] pp 1-6   Does Not Comply Corrected Diskette Needed   insufficient. Give source of genetic material (see item 11 on Error Summary Sheet)

RAW SEQUENCE LISTING   DATE: 08/07/2006
PATENT APPLICATION: US/10/587,370   TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt
Output Set: N:\CRF4\08072006\J587370.raw

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 63 <222> LOCATION: (18)..(20)
 64 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 66 <220> FEATURE:
 67 <221> NAME/KEY: MISC_FEATURE
 68 <222> LOCATION: (22)..(29)
 69 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 71 <220> FEATURE:
 72 <221> NAME/KEY: MISC_FEATURE
 73 <222> LOCATION: (31)..(33)
 74 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
 76 <400> SEQUENCE: 1
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W--> 78 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
```
 79 1               5                   10                  15
 82 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
 83             20                  25                  30
 86 Xaa Cys
 90 <210> SEQ ID NO: 2
 91 <211> LENGTH: 40
 92 <212> TYPE: PRT
 93 <213> ORGANISM: Artificial Sequence
 95 <220> FEATURE:
 96 <223> OTHER INFORMATION: Formula I
 99 <220> FEATURE:
100 <221> NAME/KEY: MISC_FEATURE
101 <222> LOCATION: (2)..(5)
102 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
104 <220> FEATURE:
105 <221> NAME/KEY: MISC_FEATURE
106 <222> LOCATION: (7)..(7)
107 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
109 <220> FEATURE:
110 <221> NAME/KEY: MISC_FEATURE
111 <222> LOCATION: (9)..(9)
112 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
114 <220> FEATURE:
115 <221> NAME/KEY: MISC_FEATURE
116 <222> LOCATION: (11)..(11)
117 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
119 <220> FEATURE:
120 <221> NAME/KEY: MISC_FEATURE
121 <222> LOCATION: (12)..(13)
122 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
124 <220> FEATURE:
125 <221> NAME/KEY: MISC_FEATURE
126 <222> LOCATION: (16)..(16)
127 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
129 <220> FEATURE:
130 <221> NAME/KEY: MISC_FEATURE
131 <222> LOCATION: (18)..(20)
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RAW SEQUENCE LISTING   DATE: 08/07/2006
PATENT APPLICATION: US/10/587,370   TIME: 09:35:48

Input Set : A:\11669.0237USWO-Sequence Listing.txt
Output Set: N:\CRF4\08072006\J587370.raw

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132 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
134 <220> FEATURE:
135 <221> NAME/KEY: MISC_FEATURE
136 <222> LOCATION: (22)..(29)
137 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
139 <220> FEATURE:
140 <221> NAME/KEY: MISC_FEATURE
141 <222> LOCATION: (31)..(33)
142 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
144 <400> SEQUENCE: 2
146 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
147 1               5                   10                  15
150 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
151             20                  25                  30
154 Xaa Cys Asn Ser Val Lys Gly Thr
155         35                  40
158 <210> SEQ ID NO: 3
159 <211> LENGTH: 34
160 <212> TYPE: PRT
161 <213> ORGANISM: Artificial Sequence
163 <220> FEATURE:
164 <223> OTHER INFORMATION: Formula I
167 <220> FEATURE:
168 <221> NAME/KEY: MISC_FEATURE
169 <222> LOCATION: (2)..(5)
170 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
172 <220> FEATURE:
173 <221> NAME/KEY: MISC_FEATURE
174 <222> LOCATION: (7)..(7)
175 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
177 <220> FEATURE:
178 <221> NAME/KEY: MISC_FEATURE
179 <222> LOCATION: (9)..(9)
180 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
182 <220> FEATURE:
183 <221> NAME/KEY: MISC_FEATURE
184 <222> LOCATION: (11)..(11)
185 <223> OTHER INFORMATION: Xaa is Leu, Val or Ile
187 <220> FEATURE:
188 <221> NAME/KEY: MISC_FEATURE
189 <222> LOCATION: (12)..(13)
190 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
192 <220> FEATURE:
193 <221> NAME/KEY: MISC_FEATURE
194 <222> LOCATION: (16)..(16)
195 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
197 <220> FEATURE:
198 <221> NAME/KEY: MISC_FEATURE
199 <222> LOCATION: (18)..(20)
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W--> 146 (marginal annotation)

```
     200 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     202 <220> FEATURE:
     203 <221> NAME/KEY: MISC_FEATURE
     204 <222> LOCATION: (22)..(29)
     205 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     207 <220> FEATURE:
     208 <221> NAME/KEY: MISC_FEATURE
     209 <222> LOCATION: (31)..(33)
     210 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     212 <400> SEQUENCE: 3
W--> 214 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
     215 1               5                   10                  15
     218 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
     219              20                  25                  30
     222 Xaa Cys
     226 <210> SEQ ID NO: 4
     227 <211> LENGTH: 34
     228 <212> TYPE: PRT
     229 <213> ORGANISM: Artificial Sequence
     231 <220> FEATURE:
     232 <223> OTHER INFORMATION: Formula I
     235 <220> FEATURE:
     236 <221> NAME/KEY: MISC_FEATURE
     237 <222> LOCATION: (2)..(5)
     238 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     240 <220> FEATURE:
     241 <221> NAME/KEY: MISC_FEATURE
     242 <222> LOCATION: (7)..(7)
     243 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     245 <220> FEATURE:
     246 <221> NAME/KEY: MISC_FEATURE
     247 <222> LOCATION: (9)..(9)
     248 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     250 <220> FEATURE:
     251 <221> NAME/KEY: MISC_FEATURE
     252 <222> LOCATION: (11)..(11)
     253 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
     255 <220> FEATURE:
     256 <221> NAME/KEY: MISC_FEATURE
     257 <222> LOCATION: (12)..(13)
     258 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     260 <220> FEATURE:
     261 <221> NAME/KEY: MISC_FEATURE
     262 <222> LOCATION: (16)..(16)
     263 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
     265 <220> FEATURE:
     266 <221> NAME/KEY: MISC_FEATURE
     267 <222> LOCATION: (18)..(18)
     268 <223> OTHER INFORMATION: Xaa is selected from the group consisting of Gln, Asp and
Ala
```

RAW SEQUENCE LISTING   DATE: 08/07/2006
PATENT APPLICATION: **US/10/587,370**   TIME: 09:35:48

Input Set : **A:\11669.0237USWO-Sequence Listing.txt**
Output Set: **N:\CRF4\08072006\J587370.raw**

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270 <220> FEATURE:
271 <221> NAME/KEY: MISC_FEATURE
272 <222> LOCATION: (19)..(20)
273 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
275 <220> FEATURE:
276 <221> NAME/KEY: MISC_FEATURE
277 <222> LOCATION: (22)..(29)
278 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
280 <220> FEATURE:
281 <221> NAME/KEY: MISC_FEATURE
282 <222> LOCATION: (31)..(33)
283 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
285 <400> SEQUENCE: 4
287 Cys Xaa Xaa Xaa Xaa Tyr Xaa Asp Xaa Leu Xaa Xaa Xaa Cys Lys Xaa
288 1               5                   10              15
291 Cys Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
292             20                  25                  30
295 Xaa Cys
299 <210> SEQ ID NO: 5
300 <211> LENGTH: 34
301 <212> TYPE: PRT
302 <213> ORGANISM: Artificial Sequence
304 <220> FEATURE:
305 <223> OTHER INFORMATION: Formula I
308 <220> FEATURE:
309 <221> NAME/KEY: MISC_FEATURE
310 <222> LOCATION: (2)..(5)
311 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
313 <220> FEATURE:
314 <221> NAME/KEY: MISC_FEATURE
315 <222> LOCATION: (7)..(7)
316 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
318 <220> FEATURE:
319 <221> NAME/KEY: MISC_FEATURE
320 <222> LOCATION: (9)..(9)
321 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
323 <220> FEATURE:
324 <221> NAME/KEY: MISC_FEATURE
325 <222> LOCATION: (11)..(11)
326 <223> OTHER INFORMATION: Xaa is any amino acid residue except Ala and cysteine
328 <220> FEATURE:
329 <221> NAME/KEY: MISC_FEATURE
330 <222> LOCATION: (12)..(13)
331 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
333 <220> FEATURE:
334 <221> NAME/KEY: MISC_FEATURE
335 <222> LOCATION: (16)..(16)
336 <223> OTHER INFORMATION: Xaa is any amino acid except cysteine
338 <220> FEATURE:
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W--> (line 287)

please correct this error in subsequent sequences.

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 08/07/2006
PATENT APPLICATION: **US/10/587,370**   TIME: 09:35:49

Input Set : **A:\11669.0237USWO-Sequence Listing.txt**
Output Set: **N:\CRF4\08072006\J587370.raw**

<u>Please Note:</u>
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:1; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,20,22,23,24,25,26,27,28,29
Seq#:1; Xaa Pos. 31,32,33
Seq#:2; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,20,22,23,24,25,26,27,28,29
Seq#:2; Xaa Pos. 31,32,33
Seq#:3; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,20,22,23,24,25,26,27,28,29
Seq#:3; Xaa Pos. 31,32,33
Seq#:4; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,20,22,23,24,25,26,27,28,29
Seq#:4; Xaa Pos. 31,32,33
Seq#:5; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,22,23,24,25,26,27,28,29,31,32
Seq#:5; Xaa Pos. 33
Seq#:10; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
Seq#:10; Xaa Pos. 28,29,31,32,33
Seq#:11; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
Seq#:11; Xaa Pos. 28,29,31,32,33
Seq#:12; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
Seq#:12; Xaa Pos. 28,29,31,32,33
Seq#:19; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
Seq#:19; Xaa Pos. 28,29,31,32,33
Seq#:28; Xaa Pos. 2,3,4,5,7,9,11,12,13,16,18,19,22,23,24,25,26,27,28,29,31
Seq#:28; Xaa Pos. 32,33
Seq#:33; Xaa Pos. 2,3,4,5,6,7,9,11,12,13,15,16,18,19,20,22,23,24,25,26,27
Seq#:33; Xaa Pos. 28,29,31,32,33

VERIFICATION SUMMARY   DATE: 08/07/2006
PATENT APPLICATION: **US/10/587,370**   TIME: 09:35:49

Input Set : **A:\11669.0237USWO-Sequence Listing.txt**
Output Set: **N:\CRF4\08072006\J587370.raw**

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number
L:78 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
M:341 Repeated in SeqNo=2
L:214 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
M:341 Repeated in SeqNo=3
L:287 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
M:341 Repeated in SeqNo=4
L:355 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
M:341 Repeated in SeqNo=5
L:548 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
M:341 Repeated in SeqNo=10
L:653 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:758 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
M:341 Repeated in SeqNo=12
L:995 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
M:341 Repeated in SeqNo=19
L:1485 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:1734 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
M:341 Repeated in SeqNo=33